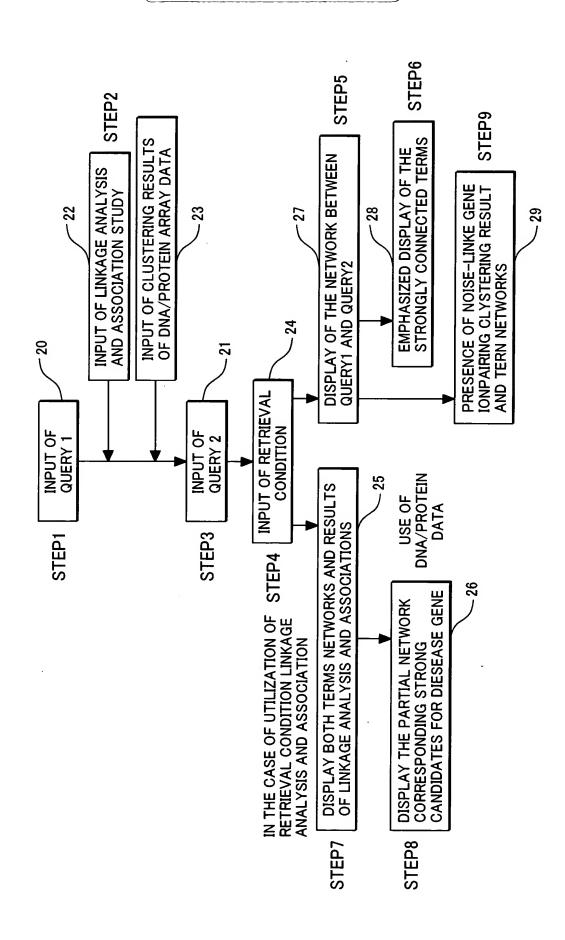
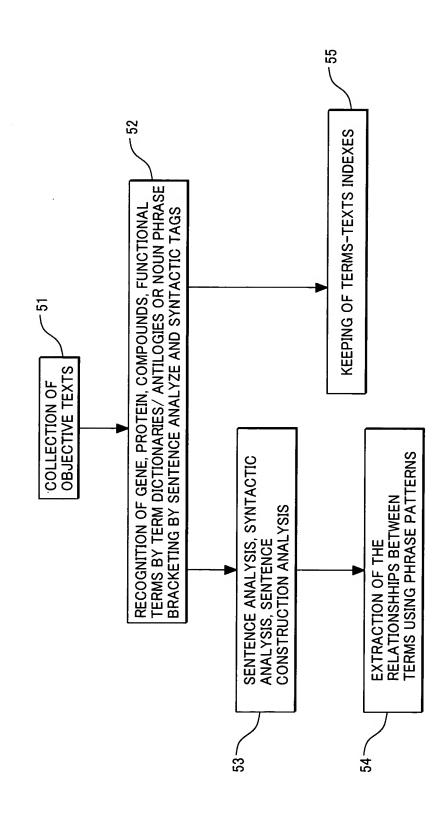


FIG.2







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PROTEIN-ID1 /CONCEPT1	PROTEIN-ID2 /CONCEPT2	RELIABILITY	EXPERIMENTAL METHOD	ORGANISM	TEXT-1D
GSC004154	GSC004160	0.95	YEAST-TWO HYBRID, MASS SPECTROMETRY	S.cerevisiae	1,2
GSC004168	GSC004154	6.0	YEAST-TWO HYBRID, MASS SPECTROMETRY	S.cerevisiae	က
CELL_DEATH.	APOPTOSIS	8.0		H.sapiens	5
GHS001223	APOPTOSIS	6:0		H.sapiens	4
GHS001223	BH2 DOMAIN	0.5		H.sapiens	9
GHS001223	GHS016577	6:0		H.sapiens	7
				H.sapiens	

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PROTEIN-ID1 /CONCEPT1	PUBMED-ID	TERM FREQUENCY	THE POSITION OF TERM IN THE TEXT (BYTE)	TEXT-ID
GSC004154	12909353	3	305,777,930	101
GSC004154	12867033	1	922	102
GSC004154	12827445	1	417	103
GSC004154	12808050	2	607,1272	104

EXPRESSION INFORMATION				NEUROBRASTOMA COT, LYMPH, BRAIN,		BLOOD, KEDNEY, SKELTAL LYMPHOCYTE,	BRAIN, SKIN	
DOMAIN INFORMATION	Ser/Thr PROTEIN KINASE DOMAIN	Ser/Thr PROTEIN KINASE DOMAIN	Ser/Thr PROTEIN KINASE DOMAIN	Ser/Thr PROTEIN KINASE DOMAIN		BH4 DOMAIN BCL-2 DOMAIN	PROTEIN 1 DOMAIN	
SEQUENCE INFORMATION	GCE000836:1e-22, GCE011584:5e-21	GCE000836:4e-32, GCE000678:1e-38	GCE000822:9e-45, GCE000667:1e-31	GCE000884:1e-164,	GSC004154:5e-21,		GCE001332:7e-95, GCE000580:8e-37	
SUBCELLULAR LOCALIZATION	CYTOPLASMIC	CYTOPLASMIC	CYTOPLASMIC	CYTOPLASMIC		MITOCONDRIAL MEMBRANE, INTRACELLULAR MEMBRANE OF THE NUCLEAR ENVELOPE, THE ENDOPLASMIC RETICULM.	INTEGRAL MEMBRANCE PROTEIN, GOLDI AND ENDOPLASMIC RETICULUM	
ORGANISM	S. cerevisiae	S. cerevisiae	S. cerevisiae	H. sapiens	C. elegans	H. sapiens	H. sapiens	•••
GENE NAME	STE11	STE20	STE7	MAPK1	K06H7.1	BCL-2	PSEN1	
OI	GSC004154	GSC004160	GSC004168	GHS012062	GCE011584	GHS001223	GHS016577	:

			_		_
EXTRACTED SENTENCE	OUR RESULTS SUGGEST THAT, IN RESPONSE TO MULTIPLE EXTRACELLULAR SIGNALS, PHOSPHORYLATION OF STE11P BY STE20P REMOVES AN AMINO-TERMINAL INHIBITORY DOMAIN, LEARNING TO ACTIVATION OF THE STE 11 PROTEIN KINASE.	STE 20P PHOSPHORYLATED STE11P ON SER302 AND/OR SER306 AND THR307 IN YEAST, RESIDUES THAT ARE CONSERVED IN MEKKS OF OTHER ORGANISMS.	INTERACTION BETWEEN STE7 AND STE11 IS BRIDGED BY STE5, SUGGESTING THE FORMATION OF A MULTIPROTEIN COMPLEX.	BCL2 PROTEINS ARE KEY MEDIATORS OF THE PROCESS OF APOPTOSIS AND LIGANDS TO THESE FAMILY OF PROTEINS HAVE BEEN DESCRIBED USING MODERN COMBINATIONAL.	
JOURNAL	CURR BIOL	CURR. BIOL.	PNAS	CURR MED CHEM	
PUBMED ID	108837245	10837245	8052657	12769779	
IMPACT FACTOR	8.4	8.4	10.8	4.9	:
TEST ID	-	2	င	4	:

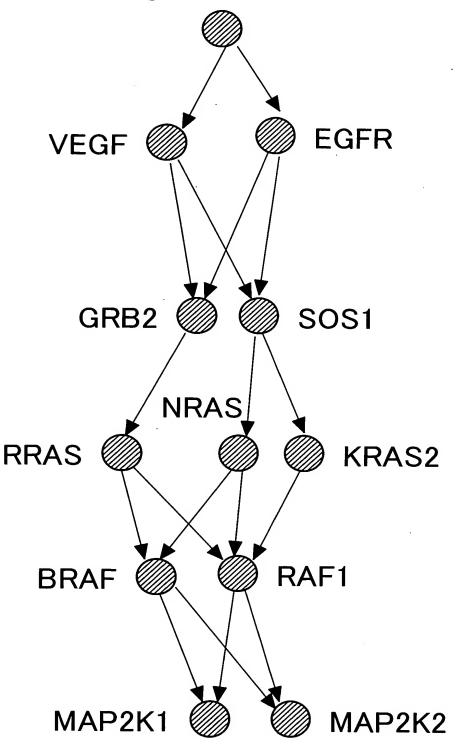
TERMS (CONCEPTS)	UPPERTERMS (CONCEPTS)
ste7	MAPK
MAP2K1	MAPK2K
MAP2KK2	MAPK2K
ste11	MAP2K
ste20	МАР3К
MAPK	ser/thr kinase
EGFR	RTK
VEGF	RTK
RRAS	RAS
NRAS	RAS
KRAS2	RAS
BRAF	RAF
RAF1	RAF
•••	

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FIG.9

growth factor

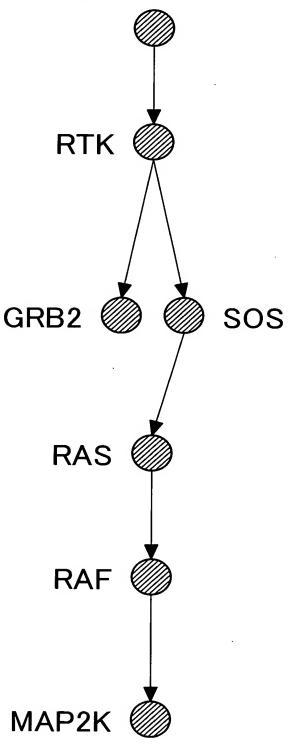


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FIG.10

growth factor



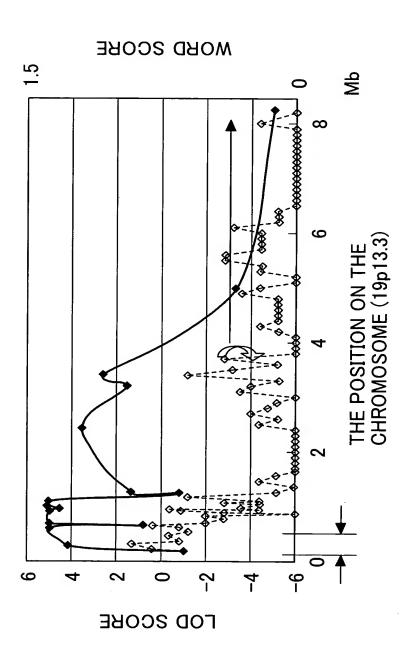
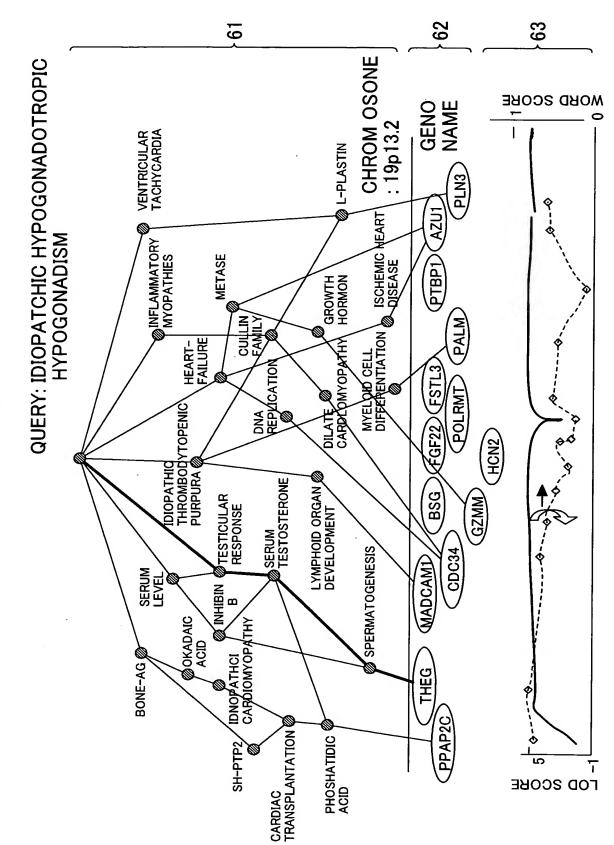


FIG.12



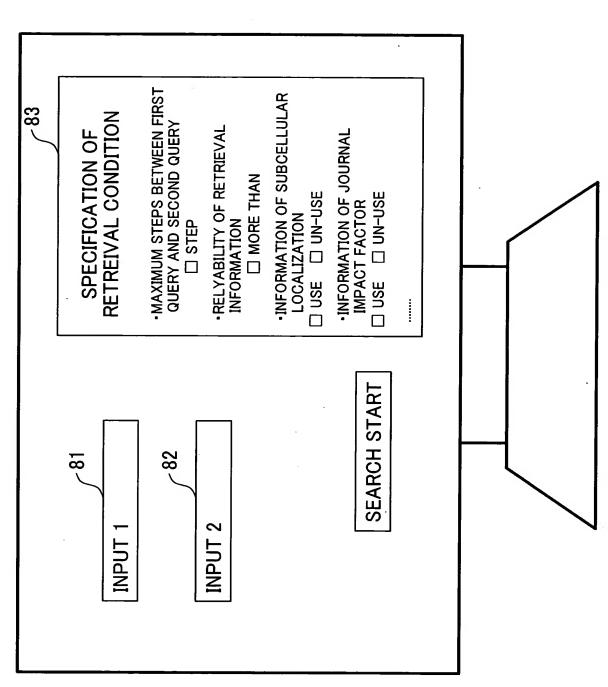
DISTANCE BETWEEN GENES

FIG.13

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**FIG.15** 

